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1 /11130	Comment							
	Serum #							

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Figure 2-1

	_
AccI 243	GT'mk_AC (0 Err) - 2 Fragment(s) 909
ApaI 402	G_GGCC'C (0 Err) - 2 Fragment(s) 750
BamHI 200	G'GATC_C (0 Err) - 2 Fragment(s) 952
EcoRI 343	G'AATT_C (0 Err) - 2 Fragment(s) 809
	A'AGCT_T (0 Err) - 2 Fragment(s) 1017
	CC's_GG (0 Err) - 2 Fragment(s) 956
	C'CATG_G (0 Err) - 4 Fragment(s) 135 288 669
	C_TGCA'G (0 Err) - 3 Fragment(s) 366 449
PvuII 570	CAG'CTG (0 Err) - 2 Fragment(s) 582
SalI 242	G'TCGA_C (0 Err) - 2 Fragment(s) 910

== Linear Map of Sequence:

1	ggggcaggcaatgagagctgcactctggctggggaaggcatgagtgacagacccacagca ccccgtccgttactctcgacgtgagaccgaccccttccgtactcactgtctgggtgtcgt	60
61	aggcggtggggtaagtgtggacctttgtgtaccagagagaacatcatggtggctttcaaa tccgccaccccattcacacctggaaacacatggtctctcttgtagtaccaccgaaagttt	120
121	HindIII description of the control	180
181	BamHI tttgttctcctcagcctgggatccaccatcaactggggtggaacagaaaagcctttaccg aaacaagaggagtcggaccctaggtggtagttgaccccaccttgtcttttcggaaatggc * * * * * * * * * * * * * * * * * * *	240

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Figure 2-2

	Acci Ncoi	
241	gtcgacatggttctcatctccctttgctttggactcagcattgcaaccatggtgcagtgc cagctgtaccaagagtagagggaaacgaaac	300
	PstI NcoI \	
301	tttggccatatcagcggtggccacatcaaccctgcagtgactgtggccatggtgtgcacc aaaccggtatagtcgccaccggtgtagttgggacgtcactgacaccgggtaccacacgtgg	360
361	aggaagatcagcatcgccaagtctgtcttctacatcgcagcccagtgcctgggggccatc tccttctagtcgtagcggttcagacagaagatgtagcgtcgggtcacggacccccggtag ^ * ^ * ^ * ^ * ^ * ^ *	420
421	attggagcaggaatcctctatctggtcacacctcccagtgtggtgggaggcctgggagtc	480
•	taacctcgtccttaggagatagaccagtgtggagggtcacaccaccctccggaccctcag	
	NcoI \	
481	accatggttcatggaaatcttaccgctggtcatggtctcctggttgagttgataatcaca tggtaccaagtacctttagaatggcgaccagtaccagaggaccaactcaactattagtgt	
	PvuII \	
541	tttcaattggtgtttactatctttgccagctgtgattccaaacggactgatgtcactggcaaagttaaccacaaatgatagaaacggtcgacactaaggtttgcctgactacagtgaccg	600
601	tcaatagctttagcaattggattttctgttgcaattggacatttattt	660
	PstI	
661	actggtgccagcatgaatcccgcccgatcctttggacctgcagttatcatgggaaattgg tgaccacggtcgtacttagggcgggctaggaaacctggacgtcaatagtaccctttaacc	720 :
	ApaI	
721	gaaaaccattggatatattgggttgggcccatcataggagctgtcctcgctggtggccttctttggtaacctatataacccaacccgggtagtatcctcgacaggagcgaccaccggaa	
	ECORI	
781	tatgagtatgtcttctgtccagatgttgaattcaaacgtcgttttaaagaagccttcagaatactcatacagaagacaggtctacaacttaagtttgcagcaaaatttcttcggaagtcg	

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Figure 2-3

											841
											901
											961
											1021
											1081
^ *	. *	* ^	•		*		*			_	1141
	NciI \ Jaccgggga tggccct * tagaagat atcttcta * ccattaag ggtaattc * attactca	NciI gacgttgaccgggga ctgcaactggccct * gtatgactagaagat catactgatctcta * cttccacccattaag gaaggtgggtaattc * atgtcatattactca	NciI description of the second of the secon	NciI gcatgtgattgacgttgaccgggga cgtacactaactgcaactggccct *	NciI tggtgcatgtgattgacgttgaccgggga accacgtacactaactgcaactggccct *	Regatgtacctccacctcctgttgtcctcagtccat *	Cettegatgtacetecacetectgttgtecteagtecat * * * * * * * * * * * * * * * * * * *	Ncil attotaaaacctggagtggtgcatgtgattgacgttgaccgggga taagattttggacctcaccacgtacactaactgcaactggccct *	NciI gacctgattctaaaacctggagtggtgcatgtgattgacgttgaccgggga ctggactaagattttggacctcaccacctactactgaccggga ctggactaagattttggacctcaccaccacgtacactaactgcaactggccct aaggggaaagaccaatctggagaggtattgtcttcagtatgactagaagat ctcccctttctggttagacctccataacagaagtcatactgatcttcta agcagacaagactccttagaactgtcctcagatttccttcc	Ncil ggatgacctgattctaaaacctggagtggtgcatgtgattgacgttgaccgggga cctactggactaagattttggacctcaccacgtacactaactgcaactggccct * * * * * * * * * * * * * * * * * * *	aaagctgcccagcaaacaaaaggaagctacatggaggtggaggacaacaggagtcaggta tttcgacgggtcgtttgttttccttcgatgtacctccactcctgttgtcctcagtccat NciI gagacggatgacctgattctaaaacctggagtggtgcatgtgattgacggtggaccccgggga ctctgcctactggactaagattttggacctcaccacgtacactaactgcaactggcccct gaggagaagaaggggaaagaccaatctggagaggtattgtcttcagtatgactagaagat ctcctcttcttcccctttctggttagacctcccataacagaagtcatactgatctcta cgcactgaaaagcagacaagactccttagaactgtcctcagatttccttcc



Citation

Algorithm Citation:

Not given.

Program Citation:

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